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WHAT WE CLAIM IS:

A DNA seguence consisting of the DNA inserts of/G-pBR322(Pst)/HFIF1, G-pBR322(Pst)/HFIF3, G-pBR322(Pst)/HFIF-6, G-pBR322(Pst)/ HFIF7, DNA sequences which hybridize to any of the foregoing DNA inserts and DNA sequences, from whatever source obtained, including natural, synthetic or semi-synthetic sources, related by mutation /including single of multiple, base substitutions, deletions, insertions and inversions

A DNA sequence according to claim 1 wherein

to any of the foregoing DNA sequences or inserts.

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DNA sequences which hybridize to any of the foregoing DNA inserts, DNA sequences, from whatever source obtained, including natural, synthetic or semi-synthetic sources, related by mutation, including single or multiple, base

said DNA sequence which hypridizes to said DNA insert is selected from the group consisting of the DNA inserts of G-pPLa-HFIF-67-12, G-pPLa/HFIF-67-12AL/9, G-pPLc-HFIF-67-8,

substitutions, deletions, insertions and inversions to any of the foregoing the sequences or inserts, and DNA sequences comprising/sequences of codons which on expression code for a polypeptide displaying similar immunological

or biological activity to a polypeptide coded for on expression of the codons of any of the foregoing DNA sequences and inserts.

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A DYA sequence according to claim 1 or 2 (3) wherein said DNA sequence which hybridizes to said DNA insert is selected from the group consisting of G-pPLa-HFIF-67-12Δ279f, G-pPLa-HFIF-67-12Δ218Ml, G-pPLa-HFIF-67-124M1, G-pHLa-HFIF-67-12419 BX-2, DNA sequences which hybridize to any of the foregoing DNA sequences, DNA sequences, from whatever source obtained, including natural, synthetic, or semi-synthetic sources, related by mutation, including single or multiple, base substitutions, delections, insertions and inversions, to any of the foregoing DNA sequences and DNA sequences comprising sequences of codons which on expression code for a polypeptide

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NOSILIEDAY PURA PA similar in immunological or biological activity to a polypeptide coded for on expression of any of the foregoing DNA sequences.

A DNA sequence according to anyone of claims 1 to 3 wherein said DNA sequence which hybridizes to said DNA insert is the DNA insert of p[325]-gAFIF-4, DNA sequences which hybridize to the foregoing DNA sequence, DNA sequences, from whatever source obtained, including natural, synthetic, or semi-synthetic sources, related by mutation, including single or multiple, base substitutions, deletions, insertions and inversions, to the foregoing DNA sequences and DNA sequence comprising sequences of codons which on expression code for a polypeptide similar in immunological or biological activity to a polypeptide coded for on expression of the foregoing DNA sequence.

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A DNA sequence according to any one of claims 2 to 4 characterized in that it is selected from the group consisting of DNA sequences of the formula: ATGACCAACAAG tgtctcctcca/aattgctctcctgttgtg/cttctccactacagctctttccatgagc TACAACTTGCTTGGATTCCTACAAAQAAGCAGCAATTTTCAGTGTCAGAAGCTCCTG TGGCAATTGAATGGGAGGCTTGAATACTGCCTCAAGAACAGGATGAACTTTGACATC CCTGAGGAGATTAAGCAGCTGCAG¢ÄGTTCCAGAAGGAGGACGCCGCATTGACCATC TATGAGATG¢TCCAGAACATCT¢TCCTATTTTCAGACAAGATTCATCTAGCACTGGC TGGAATGAGACTATTGTTGAGAACCTCCTGGCTAATGTCTATCATCAGATAAACCAT CTGAAGACAGTCCTGGAAGAAAACTGGAGAAAGAAGATTTCACCAGGGGAAAACTC ATGAGCAGTCTGCACCTGAAAAGATATTATGGGAGGATTCTGCATTACCTGAAGGCC AAGGAGTA¢AGTCACTOTGCCTGGACCATAGTCAGAGTGGAAATCCTAAGGAACTTT TACTTCATTAACAGACTTACAGGTTACCTCCGAAAC, ATGAGCTACAACTTGCTT ggattcc4acaaagaagcagcaat£ttcagtgtcagaagctcctgtggcaattgaat GGGAGGCTTGAATACTGCCTCAAGTACAGGATGAACTTTGACATCCCTGAGGAGATT aagcagcigcagcagttccagaaggaggacgccgcattgaccatctatgagatgctc CAGAACATC/TTGCTATTTTCAGA AAGATTCATCTAGCACTGGCTGGAATGAGACT ATTGTTGAGAACCTCCTGGCTAATGTCTATCATCAGATAAACCATCTGAAGACAGTC CTGGAN/AAAAACTGGAGAAAGAA GATTTCACCAGGGGAAAACTCATGAGCAGTCTG cacctsaaaagatattatgggaggattctgcattacctgaaggccaaggagtacagt CACTGTGCCTGGACCATAGTCAGAGTGGAAATCCTAAGGAACTTTTACTTCATTAAC AGACTTACAGGTTACCTCCGAAAd and derivatives thereof, said

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fragments and derivatives coding for polypeptides displaying an immunological or biological activity of IFN-β.

6. A recombinant DNA molecule characterized by a DNA sequence according to any one of the preceding claims.

7 The recombinant DNA molecule according to claim 6, characterized in that said DNA sequence is operatively linked to an expression control sequence.

8 The recombinant DNA molecule according to claim 7, characterized in that said expression control sequence is selected from the group consisting of a <u>lac</u> system, a β-lac system, a <u>tro</u> system, major operator and promotor regions of phage the control region of fd coat protein, and other sequences which control the expression of genes of prokaryotic or eukaryotic cells and their viruses.

The recombinant DNA molecule according to claim 7 or 8 selected from the group consisting of G-pPLa-HFIF-67-12, G-pPLa-HFIF-67-12A/19, and G-pPLa-HFIF-67-8.

The recombinant DNA molecule according to claim 7 or 8 characterized in that it is selected from the group consisting of G-pPIa-HFIF-67-12A279T, 6-pPLa-HFIF-67-12A218M1, 6-pPLa-HFIF-67-12AM1, and G-pPLa-HFIF-67-12A19 BX-2

11. A host transformed with at least one recombinant DNA molecule, said recombinant DNA molecule being selected from the group consisting of recombinant DNA molecules according to any one of claims 6 to 10.

The host of claim 11 selected from the group consisting of strains of <u>E. coli</u>, <u>Pseudomonas</u>, <u>Bacillus subtilis</u>, <u>Bacillus stearothermophilus</u>, other bacilli, yeasts, other fungi, mouse or other animal or plant hosts and human tissue cells.

The transformed host according to claim 11 or 12 selected from the group consisting of <u>E. coli</u> HB101 (G-pBR322(Pst)/HFIF1), <u>E. coli</u> HB101 (G-pBR322(Pst)/HFIF3), <u>E. coli</u> HB101 (G-pBR322(Pst)/HFIF6) and <u>E. coli</u> HB101 (G-pBR322(Pst)/HFIF7).

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or 12 selected from the group consisting of E.coli M5219 (G-pPLa-HFIF-67-12), H.coli K12AHI (G-pPLa-HFIF-67-12), E.coli M5219 (G-pPLa-HFIF-67-12A19), E.coli M5219 (G-pPLa-HFIF-67-8) and E.coli K12AHI (G-pPLa-HFIF-67-8)

The transformed host according to claim 11 and 12 characterized in that it is selected from the group consisting of E.coli M5219 (G-pPLa-HFIF-67-12Δ279T), E.coli M5219 (pPLa-HFIF-67-12Δ218M1), E.coli M5219 (pPLa-HFIF-67-12ΔMI), E.coli K12ΔHI (pPLa-HFIF-67-12Δ19 bX-2).

thereof displaying an immunological or biological activity of human fibroblast interferon produced by the transformed host according to any one of claims 11 to 15.

17. A polypeptide characterized in that it is coded for by a DNA sequence according to any one of claims 1

18. A polypertide or fragments and derivatives thereof according to claim 16 or 17 and being IFN-β.

19 A polymentide or/fragments and derivatives thereof according to any one of claims 16 to 18 characterized in that it is selected from the group consisting of polypeptides of the formula: /Met-Thr-Asn-Lys-Cys-Leu-Leu-Gln-Ile-Ala-Leu-Leu-Leu-Cys-The-Ser-Thr-Thr-Ala-Leu-Ser-Met-Ser-Tyr-Asn-Leu-Leu-Gly-Phe-Leu-Gln-Arg-Ser-Ser-Asn-Phe-Gln-Cys-Gln-Lys-Leu-Leu-Trp-Gln-Leu-Asn-Gly-Arg-Leu-Glu-Tyr-Cys-Leu-Lys-Asp-Ard-Met-Asn-Phe-Asp-Ile-Pro-Glu-Glu-Ile-Lys-Gln-Leu-Gln-Gln-Phe-&In-Lys-Glu-Asp-Ala-Ala-Leu-Thr-Ile-Tyr-Glu-Met-Leu-Gin-Asn-Ile-Phe-Ala-Ile-Phe-Arg-Gln-Asp-Ser-Ser-Jhr-Gly-Trp-Asn-Glu-Thr-Ile-Val-Glu-Asn-Leu-Leu-Ala-Asn-Val-Tyr-His-Gln-Ile-Asn-His-Leu-Lys-Thr-Val/Leu-Glu/Lys-Leu-Glu-Lys-Glu-Asp-Phe-Thr-Arg-Gly-Lys-Lex-Met-Ser-Ser-Leu-His-Leu-Lys-Arg-Tyr-Tyr-Gly-Arg-Ile-Leu-His-Tyf-Leu-Lys-Ala-Lys-Glu-Tyr-Ser-His-Cys-Ala-Trp-Thr-Ile-Val-Arg-Val-Glu-Ile-Leu-Arg-Asn-Phe-Tyr Phe-tle-Asn-Arg-Leu-Thr-Gly-Tyr-Leu-Arg-Asn, Met-SerCANCELLED PYA

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Tyr-Asn-Leu-Leu-Gly-Phe-Leu-Gln-Arg-Ser-Ser-Asn-Phe-Gln-Cys-Gln-Lys-Leu-Leu-Trp-Gln-Leu-Asn-Gly-Arg-Leu-Glu-Tyk Cys-Leu-Lys-Asp-Arg-Met-Asn-Phe-Asp-Ile-Pro-Glu-Glu-Vie-Lys-Gln-Leu-Gln-Gln-Phe-Gln-Lys-G/u-Asp-Ala-Ala-Leu/Thr-Ile-Tyr-Glu-Met-Leu-Gln-Asn-lle-Phe-Ala-Ile-Phe-Arg-Gln-Asp-Ser-Ser-Ser-Thr-Gly-Trp-Asp-Glu-Thr-Ile-Val-Glu-Asn-Leu-Leu-Ala-Asn-Val-Tyr-His-GIn-Ile-Asn-His-Len-Lys-Thr-Val-Leu-Glu-Glu-Lys-Leu-Glu-Lys-Glu-Asp-Phe-Thr-Arg-Gly-Lys-Leu-Met-Ser-Ser-Leu-His-Leu-Lys-Arg-Tyr-Tyr-Gly-Arg-Ile-Leu-His-Tyr-Leu-Lys-Ala-Lys-Glu-Tyr-\$er-His-Cys-Ala-Trp-Thr-Ile-Val-Arg-Val-Glu-Ile-Leu-Arg-Asn-Phe-Tyr-Phe-Ile-Asn-Arg-Leu-Thr-Gly-Tyr-Leu-Arg-Asn/ and polypeptides from whatever source obtained related to any of the foregoing polypeptides by mutation, including single or multiple, base substitutions, deletions, insertions and inversions, to any of the DNA sequences which code for them.

20. A method for producing a recombinant DNA molecule characterized by the step of introducing into a cloning vehicle a DNA sequence according to any one of claims 1-5

The method according to claim 20 characterized by the additional step of introducing into said cloning vehicle an expression control sequence according to claim 8, said expression control sequence being introduced into said cloning vehicle so as to control and to regulate the expression of said DNA sequence.

27) A method for transforming a host characterized by the step of introducing into a host a recombinant DNA molecule according to any one of claims 6 to 10.

23.) A method for producing a polypeptide displaying an immunological or biological activity of human fibroblast interferon characterized by the steps of transforming an appropriate host with a recombinant DNA molecule according to any one of claims 8 to 10; culturing said host, and collecting said polypeptide.

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20/3/81 per A 35 37 CF 1.462 ized in that the host is selected from the group consisting of strains of E. coli, Pseudomonas, Bacillus subtilis, Bacillus stearothermophilus, other bacilli, yeasts, fungi, animal or plant hosts, and human tissue cells.

25. A method for producing a polypeptide displaying an immunological or biological activity of human fibroblast interferon characterized by the steps of culturing a host transformed by a recombinant DNA molecule according to any one of claims 8 to 10 and collecting said polypeptide.

26. A process for selecting a DNA sequence coding for a polypeptide displaying an immunological or biological activity of Hulfs-β from a group of DNA sequences characterized by the step of determining which of said DNA sequences hybridizes to a DNA sequence according to any one of claims 1-5.

27. The process of claim 26 wherein said DNA sequence screened is selected from the group consisting of DNA sequences from natural sources, synthetic DNA sequences, DNA sequences from recombinant DNA molecules and DNA sequences which are a combination of any of the foregoing DNA sequences.

28. A composition for treating human viruses or treating human cancers or tumors characterized by at least one polypeptide selected from the group consisting of the polypeptides, according to any one of claims 16 to 18.

29. A composition for treating howing wiral infections characterised by at least one polypoptide selected from the group consisting of polypoptides, according to any one of claims 16 to 18 a

27.20. A method for treating human viruses or treating human cancers or tumors characterized by the step of administering to said humans in a pharmaceutically acceptable manner an effective amount of a composition according to claim 28.

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sl: A method for treating bovine viral infections characterized by the step of administering to eaid animals in a pharmacoutically asseptable manner an effective amount of a composition asserding to claim 39.

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could be by a DNA sequence that danset by briding to a DNA sequence according to any one of claims 1 to 5 and which does playe an immuno logical activity of IEN-B.

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ADDED CLAIMS 31-37 per A

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